0506



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/083,936A	
Source:	OIPE	
Date Processed by STIC:	8/7/02	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
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 - U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
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Revised 01/29/2002



Does Not Comply Corrected Diskette Needed

OIPE

DATE: 08/07/2002 RAW SEQUENCE LISTING TIME: 09:48:31 PATENT APPLICATION: US/10/083,936A

Input Set : A:\EP.txt

Output Set: N:\CRF3\08072002\J083936A.raw

3 <110> APPLICANT: Moore, Jeffrey G 5 <120> TITLE OF INVENTION: Compositions and Methods for Protecting Tissues and Cells from Damage, and For Repairing Damaged Tissues 8 <130> FILE REFERENCE: 108236.130 10 <140> CURRENT APPLICATION NUMBER: 10/083,936A

11 <141> CURRENT FILING DATE: 2002-02-27 13 <150> PRIOR APPLICATION NUMBER: US 60/271,666; US 60/302,716

W--> 14 <151> PRIOR FILING DATE: 2001-02-27; 2001-07-03

16 <160> NUMBER OF SEQ ID NOS: 10 18 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

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166 <212> TYPE: PRT
167 <213> ORGANISM: Phaseolus vulgaris
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176 Leu Thr Lys Leu Asp Ser Gly Gly Asn Pro Val Gly Ala Ser Val Gly
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179 Arg Val Leu Phe Ser Ala Pro Phe His Leu Trp Glu Asn Ser Met Ala
182 Val Ser Ser Phe Glu Thr Asn Leu Thr Ile Gln Ile Ser Thr Pro His
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 197 Asn Ala Asn Ile Asp Pro Asn Tyr Arg His Ile Gly Ile Asp Val Asn
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 203 Thr Ala Thr Ala Arg Ile Ser Tyr Asn Ser Ala Ser Lys Lys Ser Thr
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/083,936A

DATE: 08/07/2002 TIME: 09:48:31

Input Set : A:\EP.txt

Output Set: N:\CRF3\08072002\J083936A.raw

206 Val Thr Thr Phe Tyr Pro Gly Met Glu Val Val Ala Leu Ser His Asp · 195 209 Val Asp Leu His Ala Glu Leu Pro Glu Trp Val Arg Val Gly Leu Ser 220 215 212 Ala Ser Thr Gly Glu Glu Lys Gln Lys Asn Thr Ile Ile Ser Trp Ser 235 230 213 225 215 Phe Thr Ser Ser Leu Lys Asn Asn Glu Val Lys Glu Pro Lys Glu Asp 250 245 218 Met Tyr Ile Ala Asn Val Val Arg Ser Tyr Thr Trp Ile Asn Asp Val 270 265 E--> 221 Leu Ser Tyr Ile Ser Asn Lys * Met Tyr Asp Ala Leu Asn Asn Asn Asn Asn Asn Asn Stop Codon
E--> 222 275 224 His Lys Tyr Val Arg Cys Ser Thr Cys Met Leu Phe Met Lys Lys 300 295 E--> 225 290 310 <210> SEQ ID NO: 10 311 <211> LENGTH: 16 312 <212> TYPE: PRT 313 <213> ORGANISM: Sphenostylis stenocarpa 315 <400> SEQUENCE: 10 E--> 316 Ala Ala Ser Asn Asn Val Val Ala Val Glu Phe Asp Thr (Xaa Pro Asn

>See p. 3 for explanation

317 1

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/10/083,936A

DATE: 08/07/2002 TIME: 09:48:32

Input Set : A:\EP.txt

Output Set: N:\CRF3\08072002\J083936A.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of <220> to <223> is MANDATORY if n's or Xaa's are present. in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:10; Xaa Pos. 14

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/083,936A

DATE: 08/07/2002

/10/083,936A TIME: 09:48:32

Input Set : A:\EP.txt

Output Set: N:\CRF3\08072002\J083936A.raw

L:14 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD

L:221 M:342 E: Invalid Stop Code On Error, STOP CODON:*

L:222 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6

M:332 Repeated in SeqNo=6

L:316 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:10